## (19) World Intellectual Property Organization

International Bureau



# 

(43) International Publication Date 4 August 2005 (04.08.2005)

**PCT** 

### (10) International Publication Number WO 2005/071113 A1

(51) International Patent Classification7:

C12Q 1/68

(21) International Application Number:

PCT/GB2005/000052

- (22) International Filing Date: 10 January 2005 (10.01.2005)
- (25) Filing Language:

English

(26) Publication Language:

English

- (30) Priority Data:
  - 0401304.1

21 January 2004 (21.01.2004)

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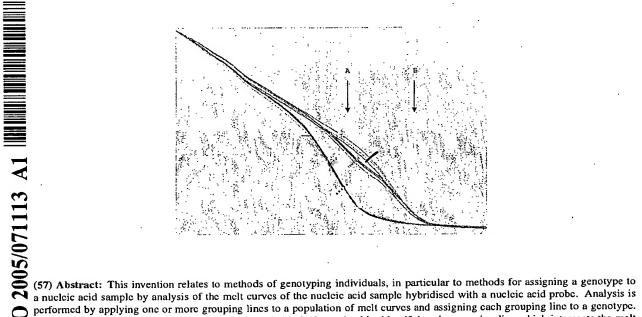
- (81) Designated States (unless otherwise indicated, for every kind of national protection available): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BW, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NA, NI, NO, NZ, OM, PG, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW.
- (84) Designated States (unless otherwise indicated, for every kind of regional protection available): ARIPO (BW, GH, GM, KE, LS, MW, MZ, NA, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IS, IT, LT, LU, MC, NL, PL, PT, RO, SE, SI, SK, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

#### Published:

with international search report

[Continued on next page]

(54) Title: METHOD OF GENOTYPING BY HYBRIDISATION ANALYSIS



performed by applying one or more grouping lines to a population of melt curves and assigning each grouping line to a genotype. The genotype of a nucleic acid sample in the population is determined by identifying the grouping line which intersects the melt curve of the nucleic acid sample.



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